

SEQUENCE LISTING

<110> USUDA, YOSHIHIRO

KURAHASHI, OSAMU

<120> METHOD FOR PRODUCING L-METHIONINE BY FERMENTATION

<130> 0010-1057-0

<140> 09/441,055

<141> 1999-11-16

<150> JP 10-326717

<151> 1998-11-17

<160> 29

<170> PatentIn version 3.1

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|---|----|
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| Met Ala Lys His Leu Phe Thr Ser Glu Ser Val Ser Glu Gly His Pro | |
| 1 5 10 15 | |

| | |
|---|----|
| gac aaa att gct gac caa att tct gat gcc gtt tta gac gcg atc ctc | 96 |
| Asp Lys Ile Ala Asp Gln Ile Ser Asp Ala Val Leu Asp Ala Ile Leu | |
| 20 25 30 | |

| | |
|---|-----|
| gaa cag gat ccg aaa gca cgc gtt gct tgc gaa acc tac gta aaa acc | 144 |
| Glu Gln Asp Pro Lys Ala Arg Val Ala Cys Glu Thr Tyr Val Lys Thr | |
| 35 40 45 | |

| | |
|---|-----|
| ggc atg gtt tta gtt ggc ggc gaa atc acc acc agc gcc tgg gta gac | 192 |
| Gly Met Val Leu Val Gly Gly Glu Ile Thr Thr Ser Ala Trp Val Asp | |
| 50 55 60 | |

| | |
|---|-----|
| atc gaa gag atc acc cgt aac acc gtt cgc gaa att ggc tat gtg cat | 240 |
| Ile Glu Glu Ile Thr Arg Asn Thr Val Arg Glu Ile Gly Tyr Val His | |
| 65 70 75 80 | |

| | |
|---|-----|
| tcc gac atg ggc ttt gac gct aac tcc tgt gcg gtt ctg agc gct atc Ser Asp Met Gly Phe Asp Ala Asn Ser Cys Ala Val Leu Ser Ala Ile 85 90 95 | 288 |
| ggc aaa cag tct cct gac atc aac cag ggc gtt gac cgt gcc gat ccg Gly Lys Gln Ser Pro Asp Ile Asn Gln Gly Val Asp Arg Ala Asp Pro 100 105 110 | 336 |
| ctg gaa cag ggc gcg ggt gac cag ggt ctg atg ttt ggc tac gca act Leu Glu Gln Gly Ala Gly Asp Gln Gly Leu Met Phe Gly Tyr Ala Thr 115 120 125 | 384 |
| aat gaa acc gac gtg ctg atg cca gca oct atc acc tat gca cac cgt Asn Glu Thr Asp Val Leu Met Pro Ala Pro Ile Thr Tyr Ala His Arg 130 135 140 | 432 |
| ctg gta cag cgt cag gct gaa gtg cgt aaa aac ggc act ctg ccg tgg Leu Val Gln Arg Gln Ala Glu Val Arg Lys Asn Gly Thr Leu Pro Trp 145 150 155 160 | 480 |
| ctg cgc ccg gac gcg aaa agc cag gtg act ttt cag tat gac gac ggc Leu Arg Pro Asp Ala Lys Ser Gln Val Thr Phe Gln Tyr Asp Asp Gly 165 170 175 | 528 |
| aaa atc gtt ggt atc gat gct gtc gtg ctt tcc act cag cac tct gaa Lys Ile Val Gly Ile Asp Ala Val Val Leu Ser Thr Gln His Ser Glu 180 185 190 | 576 |
| gag atc gac cag aaa tcg ctg caa gaa gcg gta atg gaa gag atc atc Glu Ile Asp Gln Lys Ser Leu Gln Glu Ala Val Met Glu Glu Ile Ile 195 200 205 | 624 |
| aag cca att ctg ccc gct gaa tgg ctg act tct gcc acc aaa ttc ttc Lys Pro Ile Leu Pro Ala Glu Trp Leu Thr Ser Ala Thr Lys Phe Phe 210 215 220 | 672 |
| atc aac ccg acc ggt cgt ttc gtt atc ggt ggc cca atg ggt gac tgc Ile Asn Pro Thr Gly Arg Phe Val Ile Gly Gly Pro Met Gly Asp Cys 225 230 235 240 | 720 |
| ggc ctg act ggt cgt aaa att atc gtt gat acc tac ggc ggc atg gcg Gly Leu Thr Gly Arg Lys Ile Ile Val Asp Thr Tyr Gly Gly Met Ala 245 250 255 | 768 |
| cgt cac ggt ggc ggt gca ttc tct ggt aaa gat cca tca aaa gtg gac Arg His Gly Gly Gly Ala Phe Ser Gly Lys Asp Pro Ser Lys Val Asp 816 | |

| 260 | 265 | 270 | |
|---|-----|---------|------|
| cgt tcc gca gcc tac gca gca cgt tat gtc gcg aaa aac atc gtt got | | | 864 |
| Arg Ser Ala Ala Tyr Ala Ala Arg Tyr Val Ala Lys Asn Ile Val Ala | | | |
| 275 | 280 | 285 | |
| gct ggc ctg gcc gat cgt tgt gaa att cag gtt tcc tac gca atc ggc | | | 912 |
| Ala Gly Leu Ala Asp Arg Cys Glu Ile Gln Val Ser Tyr Ala Ile Gly | | | |
| 290 | 295 | 300 | |
| gtg gct gaa ccg acc tcc atc atg gta gaa act ttc ggt act gag aaa | | | 960 |
| Val Ala Glu Pro Thr Ser Ile Met Val Glu Thr Phe Gly Thr Glu Lys | | | |
| 305 | 310 | 315 320 | |
| gtg cct tct gaa caa ctg acc ctg ctg gta cgt gag ttc ttc gac ctg | | | 1008 |
| Val Pro Ser Glu Gln Leu Thr Leu Leu Val Arg Glu Phe Phe Asp Leu | | | |
| 325 | 330 | 335 | |
| cgc cca tac ggt ctg att cag atg ctg gat ctg ctg cac ccg atc tac | | | 1056 |
| Arg Pro Tyr Gly Leu Ile Gln Met Leu Asp Leu Leu His Pro Ile Tyr | | | |
| 340 | 345 | 350 | |
| aaa gaa acc gca gca tac ggt cac ttt ggt cgt gaa cat ttc ccg tgg | | | 1104 |
| Lys Glu Thr Ala Ala Tyr Gly His Phe Gly Arg Glu His Phe Pro Trp | | | |
| 355 | 360 | 365 | |
| gaa aaa acc gac aaa gcg cag ctg ctg cgc gat gct gcc ggt ctg aag | | | 1152 |
| Glu Lys Thr Asp Lys Ala Gln Leu Leu Arg Asp Ala Ala Gly Leu Lys | | | |
| 370 | 375 | 380 | |

taa 1155

<210> 18

<211> 384

<212> PRT

<213> Escherichia coli

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Met Ala Lys His Leu Phe Thr Ser Glu Ser Val Ser Glu Gly His Pro

| | | | |
|---|-----|-----|-----|
| 1 | 5 | 10 | 15 |
| Asp Lys Ile Ala Asp Gln Ile Ser Asp Ala Val Leu Asp Ala Ile Leu | | | |
| 20 | 25 | 30 | |
| Glu Gln Asp Pro Lys Ala Arg Val Ala Cys Glu Thr Tyr Val Lys Thr | | | |
| 35 | 40 | 45 | |
| Gly Met Val Leu Val Gly Gly Glu Ile Thr Thr Ser Ala Trp Val Asp | | | |
| 50 | 55 | 60 | |
| Ile Glu Glu Ile Thr Arg Asn Thr Val Arg Glu Ile Gly Tyr Val His | | | |
| 65 | 70 | 75 | 80 |
| Ser Asp Met Gly Phe Asp Ala Asn Ser Cys Ala Val Leu Ser Ala Ile | | | |
| 85 | 90 | 95 | |
| Gly Lys Gln Ser Pro Asp Ile Asn Gln Gly Val Asp Arg Ala Asp Pro | | | |
| 100 | 105 | 110 | |
| Leu Glu Gln Gly Ala Gly Asp Gln Gly Leu Met Phe Gly Tyr Ala Thr | | | |
| 115 | 120 | 125 | |
| Asn Glu Thr Asp Val Leu Met Pro Ala Pro Ile Thr Tyr Ala His Arg | | | |
| 130 | 135 | 140 | |
| Leu Val Gln Arg Gln Ala Glu Val Arg Lys Asn Gly Thr Leu Pro Trp | | | |
| 145 | 150 | 155 | 160 |
| Leu Arg Pro Asp Ala Lys Ser Gln Val Thr Phe Gln Tyr Asp Asp Gly | | | |
| 165 | 170 | 175 | |
| Lys Ile Val Gly Ile Asp Ala Val Val Leu Ser Thr Gln His Ser Glu | | | |
| 180 | 185 | 190 | |

Glu Ile Asp Gln Lys Ser Leu Gln Glu Ala Val Met Glu Glu Ile Ile
195 200 205

Lys Pro Ile Leu Pro Ala Glu Trp Leu Thr Ser Ala Thr Lys Phe Phe
210 215 220

Ile Asn Pro Thr Gly Arg Phe Val Ile Gly Gly Pro Met Gly Asp Cys
225 230 235 240

Gly Leu Thr Gly Arg Lys Ile Ile Val Asp Thr Tyr Gly Gly Met Ala
245 250 255

Arg His Gly Gly Gly Ala Phe Ser Gly Lys Asp Pro Ser Lys Val Asp
260 265 270

Arg Ser Ala Ala Tyr Ala Ala Arg Tyr Val Ala Lys Asn Ile Val Ala
275 280 285

Ala Gly Leu Ala Asp Arg Cys Glu Ile Gln Val Ser Tyr Ala Ile Gly
290 295 300

Val Ala Glu Pro Thr Ser Ile Met Val Glu Thr Phe Gly Thr Glu Lys
305 310 315 320

Val Pro Ser Glu Gln Leu Thr Leu Leu Val Arg Glu Phe Phe Asp Leu
325 330 335

Arg Pro Tyr Gly Leu Ile Gln Met Leu Asp Leu Leu His Pro Ile Tyr
340 345 350

Lys Glu Thr Ala Ala Tyr Gly His Phe Gly Arg Glu His Phe Pro Trp
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Glu Lys Thr Asp Lys Ala Gln Leu Leu Arg Asp Ala Ala Gly Leu Lys
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28

<210> 21

<211> 28

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<213> Artificial Sequence

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gggcatgctg tagtgaggta atcaggtt

28

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28

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tgtctgctgg gcggtaca

18

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agagagtttt tcggtgcg

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<222> (1)..(927)

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48

gaa gaa aac gtc ttt gtg atg aca act tct cgt gcg tct ggt cag gaa
Glu Glu Asn Val Phe Val Met Thr Thr Ser Arg Ala Ser Gly Gln Glu
20 25 30

96

att cgt cca ctt aag gtt ctg atc ctt aac ctg atg ccg aag aag att

144

| | |
|---|-----|
| Ile Arg Pro Leu Lys Val Leu Ile Leu Asn Leu Met Pro Lys Lys Ile | |
| 35 40 45 | |
| gaa act gaa aat cag ttt ctg cgc ctg ctt tca aac tca cct ttg cag | 192 |
| Glu Thr Glu Asn Gln Phe Leu Arg Leu Leu Ser Asn Ser Pro Leu Gln | |
| 50 55 60 | |
| gtc gat att cag ctg ttg cgc atc gat tcc cgt gaa tcg cgc aac acg | 240 |
| Val Asp Ile Gln Leu Leu Arg Ile Asp Ser Arg Glu Ser Arg Asn Thr | |
| 65 70 75 80 | |
| ccc gca gag cat ctg aac aac ttc tac tgt aac ttt gaa gat att cag | 288 |
| Pro Ala Glu His Leu Asn Asn Phe Tyr Cys Asn Phe Glu Asp Ile Gln | |
| 85 90 95 | |
| gat cag aac ttt gac ggt ttg att gta act ggt gcg ccg ctg ggc ctg | 336 |
| Asp Gln Asn Phe Asp Gly Leu Ile Val Thr Gly Ala Pro Leu Gly Leu | |
| 100 105 110 | |
| gtg gag ttt aat gat gtc got tac tgg ccg cag atc aaa cag gtg ctg | 384 |
| Val Glu Phe Asn Asp Val Ala Tyr Trp Pro Gln Ile Lys Gln Val Leu | |
| 115 120 125 | |
| gag tgg tcg aaa gat cac gtc acc tcg acg ctg ttt gtc tgc tgg gcg | 432 |
| Glu Trp Ser Lys Asp His Val Thr Ser Thr Leu Phe Val Cys Trp Ala | |
| 130 135 140 | |
| gta cag gcc gcg ctc aat atc ctc tac ggc att cct aag caa act cgc | 480 |
| Val Gln Ala Ala Leu Asn Ile Leu Tyr Gly Ile Pro Lys Gln Thr Arg | |
| 145 150 155 160 | |
| acc gaa aaa ctc tct ggc gtt tac gag cat cat att ctc cat cct cat | 528 |
| Thr Glu Lys Leu Ser Gly Val Tyr Glu His His Ile Leu His Pro His | |
| 165 170 175 | |
| gcg ctt ctg acg cgt ggc ttt gat gat tca ttc ctg gca ccg cat tcg | 576 |
| Ala Leu Leu Thr Arg Gly Phe Asp Asp Ser Phe Leu Ala Pro His Ser | |
| 180 185 190 | |
| cgc tat gct gac ttt ccg gca gcg ttg att cgt gat tac acc gat ctg | 624 |
| Arg Tyr Ala Asp Phe Pro Ala Ala Leu Ile Arg Asp Tyr Thr Asp Leu | |
| 195 200 205 | |
| gaa att ctg gca gag acg gaa gaa ggg gat gca tat ctg ttt gcc agt | 672 |
| Glu Ile Leu Ala Glu Thr Glu Glu Gly Asp Ala Tyr Leu Phe Ala Ser | |
| 210 215 220 | |

aaa gat aag cgc att gcc ttt gtg acg ggc cat ccc gaa tat gat gcg 720
 Lys Asp Lys Arg Ile Ala Phe Val Thr Gly His Pro Glu Tyr Asp Ala
 225 230 235 240

caa acg ctg gcg cag gaa ttt ttc cgc gat gtg gaa gcc gga cta gac 768
 Gln Thr Leu Ala Gln Glu Phe Phe Arg Asp Val Glu Ala Gly Leu Asp
 245 250 255

ccg gat gta ccg tat aac tat ttc ccg cac aat gat ccg caa aat aca 816
 Pro Asp Val Pro Tyr Asn Tyr Phe Pro His Asn Asp Pro Gln Asn Thr
 260 265 270

ccg cga gcg agc tgg cgt agt cac ggt aat tta ctg ttt acc aac tgg 864
 Pro Arg Ala Ser Trp Arg Ser His Gly Asn Leu Leu Phe Thr Asn Trp
 275 280 285

ctc aac tat tac gtc tac cag atc acg cca tac gat cta cgg cac atg 912
 Leu Asn Tyr Tyr Val Tyr Gln Ile Thr Pro Tyr Asp Leu Arg His Met
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aat cca acg ctg gat taa 930
 Asn Pro Thr Leu Asp
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<211> 309

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<213> Escherichia coli

<400> 26

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 20 25 30

Ile Arg Pro Leu Lys Val Leu Ile Leu Asn Leu Met Pro Lys Lys Ile

35

40

45

Glu Thr Glu Asn Gln Phe Leu Arg Leu Leu Ser Asn Ser Pro Leu Gln
 50 55 60

Val Asp Ile Gln Leu Leu Arg Ile Asp Ser Arg Glu Ser Arg Asn Thr
 65 70 75 80

Pro Ala Glu His Leu Asn Asn Phe Tyr Cys Asn Phe Glu Asp Ile Gln
 85 90 95

Asp Gln Asn Phe Asp Gly Leu Ile Val Thr Gly Ala Pro Leu Gly Leu
 100 105 110

Val Glu Phe Asn Asp Val Ala Tyr Trp Pro Gln Ile Lys Gln Val Leu
 115 120 125

Glu Trp Ser Lys Asp His Val Thr Ser Thr Leu Phe Val Cys Trp Ala
 130 135 140

Val Gln Ala Ala Leu Asn Ile Leu Tyr Gly Ile Pro Lys Gln Thr Arg
 145 150 155 160

Thr Glu Lys Leu Ser Gly Val Tyr Glu His His Ile Leu His Pro His
 165 170 175

Ala Leu Leu Thr Arg Gly Phe Asp Asp Ser Phe Leu Ala Pro His Ser
 180 185 190

Arg Tyr Ala Asp Phe Pro Ala Ala Leu Ile Arg Asp Tyr Thr Asp Leu
 195 200 205

Glu Ile Leu Ala Glu Thr Glu Glu Gly Asp Ala Tyr Leu Phe Ala Ser
 210 215 220

Lys Asp Lys Arg Ile Ala Phe Val Thr Gly His Pro Glu Tyr Asp Ala
 225 230 235 240

Gln Thr Leu Ala Gln Glu Phe Phe Arg Asp Val Glu Ala Gly Leu Asp
 245 250 255

Pro Asp Val Pro Tyr Asn Tyr Phe Pro His Asn Asp Pro Gln Asn Thr
 260 265 270

Pro Arg Ala Ser Trp Arg Ser His Gly Asn Leu Leu Phe Thr Asn Trp
 275 280 285

Leu Asn Tyr Tyr Val Tyr Gln Ile Thr Pro Tyr Asp Leu Arg His Met
 290 295 300

Asn Pro Thr Leu Asp
 305

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ccagacgcac aagaagttgt c

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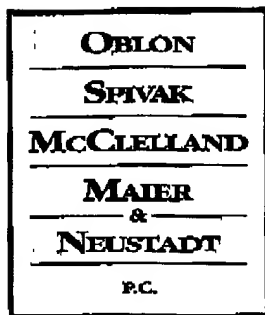
<212> PRT

<213> Escherichia coli

<400> 29

Ala Met Leu Pro Val

1 5



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| | |
|--|---|
| TO <u>Christian L. Fronda</u> <small>NAME</small> <u>U.S. Patent and Trademark Office</u> <small>COMPANY/FIRM</small> NUMBER OF PAGES INCLUDING COVER: <u>21</u> | <u>May 23, 2002</u> <small>DATE</small> <u>703-746-5036</u> <i>2nd DAY</i> <small>FAX #</small> CONFIRM FAX: <input type="checkbox"/> YES <input type="checkbox"/> NO |
| FROM <u>Thomas Barnes</u> <small>NAME</small> <u>703-412-3525</u> <small>DIRECT PHONE #</small> | <u>0010-1057-0</u> <small>OUR REFERENCE</small> <u>Serial No. 09/441,055</u> <small>YOUR REFERENCE</small> |

MESSAGE

In accordance with your instructions in our telephone discussion of May 22, 2002, I have attached a copy of the Sequence Listing filed on February 19, 2002.

Please call if you have any questions.

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